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OM protein - protein search, using sw model

August 28, 2003, 18:21:02 ; Search time 37.0909 Seconds (without alignments) 51.353 Million cell updates/sec US-09-743-225-8 58

Run on:

BLOSUM62 Gapop 10.0 , Gapext 0.5 1 NTLKTPRVGGXA 12 Title: Perfect score: Sequence: Scoring table:

1107863 seqs, 158726573 residues Searched:

1107863 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

/SIDS1/gcgdata/geneseg/genesegp-embl/AA1984.1 /SIDS1/gcgdata/geneseg/genesegp-embl/AA1985.1 /SIDS1/gcgdata/geneseg/genesegp-embl/AA1986.1 // SIDSI/gcgdata/geneseq/geneseqp-embl/AA1991/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1991/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1991/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1993/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1994/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1994/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1994/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1994/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1904/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1904/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1904/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1904/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1904/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1904/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1904/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1904/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1904/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1904/SIDSI/gcgdata/geneseq/geneseqp-embl/AA1904/SIDSI/gcgdata/geneseq/geneseq/geneseqp-embl/AA1904/SIDSI/gcgdata/geneseq/geneseq/geneseqp-embl/AA1904/SIDSI/gcgdata/geneseq/geneseq/geneseq/geneseqp-embl/AA1904/SIDSI/gcgdata/geneseq/gen /gcgdata/geneseq/geneseqp-emb1/AA1980. /SIDS1 /SIDS1, /SIDS1

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution. /SIDS1/gcgdata/geneseq/geneseqp-emb1/AA2001.DAT /SIDS1/gcgdata/geneseq/geneseqp-emb1/AA2002.DAT /SIDS1/gcgdata/geneseq/geneseqp-emb1/AA2003.DAT

## SUMMARIES

	Description	Monopeptide which	Peptide which inhi	Beta-2GPI Ab bindi	Exemplary pharmaco	Beta-2GPI Ab bindi	Exemplary pharmaco	N. gonorrhoeae ami	Human 5' EST secre	Human 5' EST secre
	ID	AAY69259	AAY69268	AAB17990	ABB73361	AAB17989	ABB73360	ABP77111	AAY11550	AAY11742
	DB	12	71	21	23	21	23	74	20	20
	Query Match Length DB ID	12	10	11	11	œ	œ	372	68	83
ø	Query Match	56 96.6	91.4	91.4	91.4	70.7	70.7	65.5	63.8	63.8
	Score	56	53	53	23	41	41	38	37 .	3.7
	Result No.	7	ď	m	4	ស	Q	7	80	6

Bifidobacterium lo	Propionibacterium	Human polypeptide,	Lactococcus lactis	•	•	Propionibacterium	D)	AMEPV fifth RNA po	<b>a</b> )	ø.				Arabidopsis thalla	Mutant PDG encoded	Mutant PDG encoded		Arabidopsis thalia		. Mutant PDG encoded			Δ.	E. coli phosphogly	Wild-type phosphog	E. coli cellular p	PDG	Mutant PDG encoded	a	m	တ	c	human	Novel human diagno	Human polypeptide
6	1	0	0	3	7	0	4	m	۳	œ	ų	7	4	Q	ι.	4	<u></u>	Ŋ	-	<b>0</b> 0	o	Ċ,	E,	Ci.	ē	80	ō,	7	بو.	2		<u>6</u>	7	7	4
ABP6629	AAU4893	AAM9375	ABB5446	AAG0639	AAG0639	AAU4196	AAB2385	ABB7763	AAM8484	AAM86368	AAM4945	ABG0887	AAG14804	AAG4846	<b>AAR3921</b>	AAR3921	AAG1480	AAG4846	AAR3922	AAR39218	AAR39220	AAR3922	<b>AAR3921</b>	AAR3921	AAR5009	AAU34698	AAR3921	AAR39217	<b>AAR39216</b>	AAG14802	AAG4.846	ABG2248	ABG3008	ABG0415	AA00168
23	22	22	23	21	21	22	21	. 23	22	22	23	22	71	71	14	14	21	21	14	14	14	14	14	14	15	22	14	14	14	71	21	22	7	22	22
388	92	140	256	319	323	330	1002	1301	28	144	200	234	332	332	364	365	367	367	374	395	403	404	405	410	410	410	411	412	414	456	456	884	884	1072	84
63.8	62.1	62.1	62.1	62.1	62.1	62.1	62.1	62.1	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	60.3	58.6
37	36	36	36	36	36	36	36	36	35	35	32	35	32	35	35	32	35	35	35	32	35	35	32	35	35	35	35	35	32	35	32	32	32	32	34
10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	56	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45

## ALIGNMENTS

Monopeptide which inhibits anti-beta-2-glycoprotein 1 antibodies. Anti-beta-2-glycoprotein l antibody; anti-B2GPI antibody; anti-phospholipid syndrome; anti-phospholipid antibody; pregnancy complication; thrombosis; coagulation dysregulation. Location/Qualifiers 11 /note= "FmocLys(Fmoc)-OH" AAY69259 standard; peptide; 12 AA. 30-MAY-2000 (first entry) Key Modified-site Synthetic. AAY69259; RESULT 1

99WO-IL00366 WO200001729-A2. 06-JUL-1999; 13-JAN-2000.

Blank M, Cabilly S, Shoenfeld Y, Katchalski-Katzir E; (YEDA ) YEDA RES & DEV CO LTD

98IL-0125262.

07-JUL-1998;

complications, thrombosis or coagulation dysregulation.

10 AA;

Sequence

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of inhibiting the biological activity of anti-beta-2-glycoprotein 1 (B2GPI) monoclonal antibodies in vitro and of inhibiting in vivo induction of experimental anti-phospholipid syndrome in mice by anti-B2GPI monoclonal antibodies. The peptides are used for diagnosis and treatment of anti-phospholipid syndrome. They may also be used for the diagnosis of anti-phospholipid antibodies with different pathogenic blofunctions which may correlate with either pregnancy
                                                                                                                                                                                                                                                                                                                                                              The present sequence represents a synthetic peptide which is capable of inhibiting the biological activity of anti-beta-2-glycoprotein 1 (B2GPI) monoclonal antibodies in vitro and of inhibiting in vivo induction of experimental anti-phospholipid syndrome in mice by anti-B2GPI monoclonal anti-phospholipid syndrome in mice by anti-B2GPI monoclonal anti-phospholipid syndrome. They may also be used for the diagnosis of anti-phospholipid antibodies with different pathogenic biofunctions which may correlate with either pregnancy complications, thrombosis or coagulation dysregulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                     Novel synthetic peptides that inhibit anti-beta-2-glycoprotein 1 antibodies, useful for diagnosis and treatment of anti-phospholipid syndrome in humans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel synthetic peptides that inhibit anti-beta-2-glycoprotein lantibodies, useful for diagnosis and treatment of anti-phospholipid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptide which inhibits anti-beta-2-glycoprotein 1 antibodies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 56;
Pred. No.
                                                                                                                                                                                                                                                                             Disclosure; Page 13; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY69268 standard; peptide; 10 AA.
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Best Local Similarity luv...
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WPI; 2000-182105/16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12 AA;
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The present invention describes composition of matter (I) comprising an Fe domain, pharmacologically active peptides, and linkers. Where (I) is: (X1)a-F1-(X2)b, where: F1 = an Fe domain; X1 and X2 = are each independently selected from -(L1)c-F1-(L1)c-F2, -(L1)c-F2, -(L4)f-F4, -(L1)c-F1-(L2)d-F2, -(L4)f-F4, -(L1)c-F1-(L2)d-F2, -(L4)f-F4, -(L1)c-F1-(L2)d-F2-(L3)e-F3-(L4)f-F4, where P1, P2, P3, and P4 = are each independently sequences of pharmacologically active peptides; L1, L2, L3, and L4 = are each independently linkers; and a, b, c, d, e, and f = are each independently of or 1, provided that at least 1 of a and b is 1. The composition can have cytostatic, antiasthmatic, thrombolytic and immunosuppressive activities. DNAs, vectors and host cells from the present invention can be used for producing pharmacoutical compositions. The compositions are useful for treating cancer, asthma, thrombosis, or autoimmune diseases. The use of an Fc domain (rather than a Fab domain) can provide a longer half-life or incorporate functions such as Fc receptor binding, protein A binding, complement fixation, and possibly placental transfer. AAA66443 to AAA69526 and AAB18003 represent uncleotide and amino acid sequences used in the exemplification of the present invention.
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                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            autoimmune disease; cytostatic; antiasthmatic; thrombolytic; VEGF; immunosuppressive; EPO; TPO; CTLA4; mimetic; IL-1; TNF; antagonist; MMF; inhibitor; erythropoietin; thrombopoietin; interleukin 1; cytotoxic T cell lymphocyte antigen 4; tumour necrosis factor; vascular endothellal growth factor; matrix metalloproteinase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel composition of matter comprising an Fc domain and pharmacologically active peptides, useful for treating cancer and
                                                                        ö
                            Length 10;
                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Modified peptide; therapeutic agent; fusion; Fc domain;
                                                                                                                                                                                                                                                                                                                                                                                                             Beta-2GPI Ab binding peptide sequence SEQ ID NO:1102.
                         Score 53; DB 21;
Pred. No. 0.0021;
                                                                      Mismatches
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                                                                                                                                                                                                                                                                     AAB17990 standard; Peptide; 11
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                         91.4%; ; 100.0%;
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                                                                      Conservative
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Ouery Match
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                                                                                                                  1 NTLKTPRVGG
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                                                                                                                                                                                                                                                                                                                    AAB17990;
                                                                                                                                                                                                                                RESULT 3
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11 AA;

Sequence

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Gaps

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multimers. (I) can have antiinflammatory, antitumour, immunosuppressive, cytostatic, antirhumatic, antiinflammatory, antifumour, immunosuppressive, cytostatic, antirhumatic, antifumour, immunosuppressive, containment, antianment, operatic, operatic, operatic, antiinfertility, haemostatic, dermatological, neuroprotective activities. (I) can be used as a therapeutic or prophylactic agent as well as for screening purposes. (I) is useful for diagnosing diseases characterised by dysfunction of their associated protein of interest, for identifying normal or abnormal proteins of interest, as a part of diagnostic kit to detect the presence of their correcting of interest in a biological sample. Additionally, (I) is useful for treating inflammatory and autoimmune diseases, tumour growth, cancer, rheumatoid arthritis, diabetic retinopathy, obesity, sleep disorders, inferitility, and neurological degenerative diseases. (I), comprising the red blood cell levels such as anaemia. The TPO-mimetic comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                    Modified peptide; mimetic; Fc domain; fusion; immunoglobulin G; IgG; EPO; erythropotetin; TPO; tumour necrosis factor alpha inhibitor; TWF-alpha inhibitor; interleukin i antagonist; IL-1 antagonist; TWF-alpha inhibitor; interleukin i antagonist; IL-1 antagonist; TWF mimetic peptide; EPO mimetic peptide; EMP; VEGF antagonist; MMP inhibitor; antiinflammatory; antitumour; immunosuppressive; antiantheumatic; antituffertility; nemostatic; dermatological; antianaemic; anorectic; antiinfertility; haemostatic; dermatological; encroprotective; inflammatory disease; autoimmune disease; tumour growth; ancer; rheumatorid arthritis; diabetic retinoptaty; infertility; obesity; sleep disorder; neurological degenerative disease; aneamia; thrombocytopaenia; metastatic tumour; systemic lupus erythematosus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   compounds are useful for treating conditions that involve an existing megakaryocyte/platelet deficiency or an expected megakaryocyte/platelet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present invention describes a vehicle-peptide molecule (I) or its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel vehicle-peptide molecule or its multimers useful for treating inflammatory and autoimmune diseases, cancer, rheumatoid arthritis, diabetic retinopathy, obesity, sleep disorders and infertility
                                             ö
  Length 11;
                                                                                                                                                                                                                                                                                                                                                                                              Exemplary pharmacologically active peptide SEQ ID NO:1100
91.4%; Score 53; DB 21; Length 11
100.0%; Pred. No. 0.0023;
.tve 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Boone TC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 39; Page 62; 176pp; English.
                                                                                                                                                                                                                                                       ABB73361 standard; Peptide; 11 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Liu C, Cheetham JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-MAY-2001; 2001WO-US14310.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-MAY-2000; 2000US-0563286
                                                                                                                                                                                                                                                                                                                                                 05-APR-2002 (first entry)
  Query Match 91.4
Best Local Similarity 100.
Matches 10; Conservative
                                                                                                                      1 NTLKTPRVGG 10
                                                                                          1 NTLKTPRVGG 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-130313/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fancon1's syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (AMGE-) AMGEN INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Feige U,
                                                                                                                                                                                                                                                                                                    ABB73361;
                                                                                                                                                                                                       RESULT 4
ABB73361
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For domain, pharmacologically active peptides, and indices. Where (I) is:

(XI)a-FI-(XZ)b, where: FI = an Fc domain; XI and X2 = are each independently selected from -(Lio-PI, -(Li)c-PI-(Li2)d-P2, (Li)c-P2, (Li)c-P1, -(Li)c-P1-(Li2)d-P2, (Li)c-P2, (Li)c-P1, P2, P3, and P4 = are each independently sequences of pharmacologically active peptides; Li, Li, Li, Li, and L4 = are each independently inkers; and a, b, c, d, e, and f = are each independently linkers; and a, b, c, d, e, and f = are each independently interest and a, b, c, d, e, and f = are each independently have cytostatic, antiasthmatic, thrombolytic and immunosuppressive activities. DNAs, vectors and host cells from the present invention can be used for producing pharmaceutical compositions. The compositions are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            an
18:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The use of an Fc domain (rather than a Fab domain) can provide a longer half-life or incorporate functions such as Fc receptor binding, protein
deficiency, such as thrombocytopaenia, aplastic anaemia, metastatic tumour which result in thrombocytopaenia, systemic lupus erythematosus, and Fanconi's syndrome. ABB72403 to ABB73426 and ABL35695 to ABL35777 represent amino acid and nucleic acid sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer, asthma, thrombosis, or autoimmune diseases
                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes composition of matter (I) comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            autoimmune disease, cytostatic; antiasthmatic; thrombolytic; VEGF; immunosuppressive; EPO; TPO; CTLA4; mimetic; IL-1; TNF; antagonist; MMF; inhibitor; erythropoietin; thrombopoietin; interleukin 1; cytotoxic T cell lymphocyte antigen 4; tumour necrosis factor; vascular endothelial growth factor; matrix metalloproteinase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel composition of matter comprising an Fc domain and pharmacologically active peptides, useful for treating cancer and autoimmune diseases
                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Modified peptide; therapeutic agent; fusion; Fc domain; cancer;
                                                                                                                                                      91.4%; Score 53; DB 23; Length 11; 100.0%; Pred. No. 0.0023; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Beta-2GPI Ab binding peptide sequence SEQ ID NO:1101.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Feige U, Liu C, Cheetham J, Boone TC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vascular endothelial growin ractor; asthma; thrombosis; pharmaceutical.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 39; Page 599; 608pp; English.
                                                                                                                                                                                                                                                                                                                                                                       AAB17989 standard; Peptide; 8 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99WO-US25044.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-OCT-2000 (first entry)
                                                                                                                                                                                              10; Conservative
                                                                                                                                                                                                                                  1 NTLKTPRVGG 10
                                                                                                                                                                                                                                                          1 NTLKTPRVGG 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 useful for treating
                                                                                                                                                                              Best Local Similarity
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                                                                                                                     11 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                               AAB17989;
                                                                                                                       Sequence
                                                                                                                                                        Query Match
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EPO; erythropoletin; TPO; tumour necrosis factor alpha inhibitor; TNF-alpha inhibitor; interleukin 1 antagonist; IL-1 antagonist; TMP; TPO mimetic peptide; EMP; VEGF antagonist; TMP; MMP inhibitor; antiinflammatory; antitumour; immunosuppressive; cytostatic; antirheumatic; antiarthritic; antidiabetic; ophthalmological; antianaemic; anorectic; antinfertility; haemostatic; dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     neuroprotective; inflammatory disease; autoimmune disease; tumour growth; cancer; rheumatoid arthritis; diabetic retinopathy; infertility; obesity; sleep disorder; neurological degenerative disease; anaemia; thrombocytopaenia; metastatic tumour; systemic lupus erythematosus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a vehicle-peptide molecule (I) or its multimers. (I) can have antiinflammatory, antitumour, immunosuppressive, cytostatic, antitheumatic, antiathritic, antidabetic, obthahimological, antianaemic, anorectic, antiminertility, heemostatic, dermatological and neuroprotective activities. (I) can be used as a therapeut or prophylactic agent as well as for screening purposes. (I) is useful for diagnosing diseases characterised by dysfunction of their associated protein of interest, for identifying normal or abnormal proteins of interest, as a part of diagnostic kit to detect the presence of their proteins of interest and autolomina diseases, tumour growth, cancer, remanatoid arthritis, diabetic retinopathy, obesity, sleep disorders, infertility, and neurological degenerative diseases. (I), comprising
A binding, complement fixation, and possibly placental transfer. AAA69443 to AAA69526 and AAB16955 to AAB18003 represent nucleotide and amino acid sequences used in the exemplification of the present invention.
                                                                                                                                                     Gapa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          peptide; mimetic; Fc domain; fusion; immunoglobulin G; IgG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel vehicle-peptide molecule or its multimers useful for treating inflammatory and autoimmune diseases, cancer, rheumatoid arthritis, diabetic retinopathy, obesity, sleep disorders and infertility -
                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                    Exemplary pharmacologically active peptide SEQ ID NO:1099.
                                                                                                                Length 8;
                                                                                                                                                   Indels
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                                                                                                              Score 41; DB 21; I
Pred. No. 9.3e+05;
0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 39; Page 62; 176pp; English.
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                                                                                                                                                                                                                                                                                                                    ABB73360 standard; Peptide; 8
                                                                                                              70.7%; 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                            Query Match
Best Local Similarity 100.0
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                                                                                                                                                                                                          1 NTLKTPRV 8
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                                                                                                                                                                                    1 NTLKTPRV
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                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Modified
                                                                                                                                                                                                                                                                                                      ABB73360
                                                                                                                                                                                                                                                                                   RESULT
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EPO-mimetic compounds are useful for treating disorders characterised by low red blood cell levels such as anaemia. The TPO-mimetic comprising compounds are useful for treating conditions that involve an existing megakaryocyte/platelet deficiency or an expected megakaryocyte/platelet deficiency or an expected megakaryocyte/platelet tumour which result in thrombocytopaenia, aplastic anaemia, metastatic tumour which result in thrombocytopaenia; systemic lupus erythematosus, and Fanconi's syndrome. ABB72403 to ABB73426 and ABL35695 to ABL35777 represent amino acid and nucleic acid sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Also disclosed are the nucleic acid molecules encoding the proteins and antibodies that specifically bind to the proteins. The composition comprising the protein, nucleic acid or antibody is useful for the manufacture of a medicament for treating or preventing N. gonorrhoeae infection, this may be in the form of a vaccine or gene therapy. Sequences given in records ABP76736-ABP81046 represent nucleic acid molecules of the invention.
                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New protein from Neisseria gonorrheae, useful for the manufacture of medicament for treating or preventing N. gonorrheae infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to proteins from Neisseria gonorrhoeae
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                                                                                                                                                                                            Length 8;
                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                          70.7%; Score 41; DB 23; I 100.0%; Pred. No. 9.3e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antibacterial; infection; vaccine; gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N. gonorrhoeae amino acid sequence SEQ ID 752.
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2; Mismatches
                                                                                                                                                                                                                        0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 244; 815pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Masignani V,
                                                                                                                                                                                                                                                                                                                                                                       ABP77111 standard; Protein; 372 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          65.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-FEB-2002; 2002WO-IB02069.
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                                                                                                                                                                       Juery Match
Bost Local Similarity 100.00
The Bost Local Similarity 100.00
                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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Best Local Similarity 70.0
Matches 7; Conservative
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182 DTLRHPRVGG 191
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Neisseria gonorrhoeae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fontana MR, Pizza M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-058415/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CHIR-) CHIRON SPA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            372 AA;
                                                                                                                                                                                                                                                                            1 NTLKTPRV
                                                                                                                                                                                                                                                          1 NTLKIPRV
                                                                                                                                                          8 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       07-MAR-2003
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                                                                                                                                                            Sequence
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                                                                                                                                                                                                                                                                                                                                       RESULT 7
ABP77111
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Human; secreted protein; EST; expressed sequence tag; diagnosis; forensic; gene therapy; chromosome mapping; signal peptide; prostate; upstream regulatory sequence; cytokine activity; cell proliferation; differentiation; haematopolesis regulation; tissue growth regulation; reproductive hormone regulation; chemotactic; chemokinetic; haemostatic; thrombolytic; anti-inflammatory; tumour inhibition.

Human 5' EST secreted protein SEQ ID No: 342

18-JUN-1999 (first entry)

AAY11742 standard; Protein; 82 AA.

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                                                                                                                                                                                                                     Human; secreted protein; EST; expressed sequence tag; diagnosis; forensic; gene therapy; chromosome mapping; signal peptide; upstream regulatory sequence; cytokine activity; cell proliferation; differentiation; haematopoiesis regulation; tissue growth regulation; reproductive hormone regulation; chemotactic; chemokinetic; haemostatic; thrombolytic; anti-inflammatory; tumour inhibition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX40251 to AAX40397 represent 5' expressed sequence tags (ESTs) for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polypeptide into a membrane, or importing a polypeptide into a cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New nucleic acids encoding human secreted proteins - obtained from cDNA libraries derived from liver, lung, large intestine, colon,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; DB 20; Length 68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dumas Milne Edwards J, Lacroix B;
                                                                                                                                                                 secreted protein SEQ ID NO:202.
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Pred. No. 17;
1; Mismatches
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AAY11550 standard; Protein; 68 AA.
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63.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98WO-IB01233.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   thyroid and pancreas tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97US-0904468
                                                                                                               (first entry)
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Best Local Similarity 63.6
Matches 7; Conservative
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N-PSDB; AAX40268.
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                                                                                                                                                                                                                                                                                                                                                                                                                          Homo saptens.
                                                                                                                                                                    5' EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              W09906439-A2
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                                                                                                            16-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-FEB-1999
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                                                       AAX11550;
                                                                                                                                                                      Human
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New isolated prostate-derived nucleic acids - used to develop products which may have cytokine, immune regulatory, haematopolesis regulating, anti-inflammatory or tumour inhibition activity

Duclert A, Dumas Milne Edwards J, Lacroix B;

(GEST ) GENSET

WPI; 1999-153780/13.

N-PSDB; AAX40464

98WO-IB01232.

31-JUL-1998; 11-FEB-1999,

WO9906550-A2. Homo sapiens

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        can be used in forensic, gene therapy and chromosome mapping procedures. The sequences can also be used for obtaining corresponding promoter asquences. The nucleic acids encoding the signal peptides can be used for directing extracellular secretion of a polypeptide or the insertion of a polypeptide into a membrane, or importing a polypeptide into a cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAX40438 to AAX40715 represent 5' expressed sequence tags (ESTs) for human secreted proteins expressed in prostate, and encode the proteins given in AAX11716 to AAX11993 respectively. The proteins given represent the signal peptide and an N-terminal fragment of a secreted protein. The nucleic acid sequences can be used for producing secreted human gene products. They can also be used for producing secreted human gene therapy. The proteins obtained may have cytokine activity, cell proliferation and differentiation activity, haematopoiesis regulating activity, tissue growth regulating activity, haematopoiesis regulating activity, receptor/ligand activity, haemostatic and thrombolytic activity, receptor/ligand activity, anti-inflammatory activity, tumour inhibition activity or other activity, and products
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 82;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 37; DB 20;
Pred. No. 20;
1; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 34; Page 517-518; 675pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABP66299 standard; Protein; 388 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            63.8%;
63.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 63.00
7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9 TLSTERIGGAA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 TLKTPRVGGXA 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  82 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABP66299;
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Gaps

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3; Indels

| | | | :|| | | TLSTERIGGAA 19 2 TLKTPRVGGXA 12

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RESULT 9 AAY11742

SAPHO syndrome; synovitis; acne; pustulosis; hypertosis; osteomyelitis; uveltis; endophthalmitis; bone; joint; central nervous system; ELISA; inflammatory lesion; acne vulgaris; ensyme linked immunosorbent assay; dermatological; osteopathic; neuroprotectant.

Propionibacterium acnes

WO200181581-A2

01-NOV-2001

Propionibacterium acnes immunogenic protein #9827.

(first entry)

27-FEB-2002

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The present invention describes a polynucleotide (I) comprising a sequence of a Bifidobacterium genome selected from the nucleotide sequence given in AB081842 and AB081843, or a sequence exhibiting at sequence 9 given in AB081843. Also described is a polynucleotide (II) encoding a fail on protein, comprising a sequence selected from 1097 sequences given in AB965258 to AB966354 ligated in frame to a polynucleotide (II) encoding a heterologous polypeptide. (I) has antidiarrheic and antidiaterial activities, and can be used as an inhibitor of Salmonella. (I) (which is a probe) is useful for the detection and/or identification of Bifidobacterium longum in a biological sample. A carrier containing the lactic acid bacterium Blidobacterium longum NGC2705 (CNCM I-2618) can be used for preventing and/or treating diarrhoea brought about by pathogenic bacteria and/or rotavirus. The carrier is a food composition selected from malk, rogunt, curd, cheese, fermented milks, milk based fermented products, ice-creams, fermented cereal based products, infant formula, pet food or a pharmaceutical composition selected from tablets, liquid bacterial suspensions, dried or all carrier containing supplement, wet oral supplement, dry tube feeding or wet tube feeding.

(I) is useful in DNA arrays or chips to carry out analysis of the expension of the Bifidobacterium gene AB081884 to AB081880 represent at the factor of the back and analysis of the seriams.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N.B. The sequence data for this patent is not represented in the printed specification but is based on sequence information supplied by the European Patent Office.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bifidobacterium related nucleotide sequences given in the Sequence
Listing from the present, invention but not mentioned further within the
                                                                                                                antidiarrheic; antibacterial; inhibitor of Salmonella; detection; identification; lactic acid bacterium; diarrhoea; pathogenic bacteria; rotavirus; food composition; pharmaceutical composition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 comprising Bifidobacterium genome sequence useful for detecting and/or identifying Bifidobacterium il sample .
                                            Bifidobacterium longum NCC2705 ORF amino acid sequence SEQ ID NO:1043.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                        Bifidobacterium longum NCC2705; Bifidobacterium; bacterial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 37; DB 23; Length 388;
Pred. No. 1e+02;
3; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; SEQ ID 1043; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel polynucleotide comprising as a probe or primer for detectiongum in a biological sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   63.8%;
60.0%;
                                                                                                                                                                                                                                                                                                                                         30-JAN-2001; 2001EP-0102050
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19-NOV-2002 (first entry)
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Matches 6; Conservative
                                                                                                                                                                                                    Bifidobacterium longum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-668397/72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     388 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            specification.
                                                                                                                                                                                                                                                  EP1227152-A1
                                                                                                                                                                                                                                                                                          31-JUL-2002.
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Propionibacterium acnes polypeptides and nucleic acids useful for vaccinating against and diagnosing infections, especially useful for

Example 1; SEQ ID No 10126; 1069pp; English.

treating acne vulgaris

Bhatia A;

Mitcham JL, Wang SS, , Jen S, Carter D;

Skeiky YAW, Persing DH, M L'maisonneuve J, Zhang Y,

(CORI-) CORIXA CORP.

WPI; 2001-616774/71. N-PSDB; AAS59543.

21-APR-2000; 2000US-199047P. 02-JUN-2000; 2000US-208841P. 07-JUL-2000; 2000US-216747P.

20-APR-2001; 2001WO-US12865

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Sequences AAU39105-AAU68017 represent Propionibacterium acnes immunogenic polypeptides. The proteins and their associated DNA sequences are used in the treatment, prevention and diagnosis of medical conditions caused by 2. Acnes. The disorders include SAPHO syndrome (synovitis, acne by pustulosis, hypertosis and osteomyelitis), uveitis and endophthalmitis. P. acnes is also involved in infections of bone, joints and the central nervous system, however it is particularly involved in the inflammatory lesions associated with acne vulgaris. A method for detecting the presence or absence of P. acnes in a patient comprises contacting a sample with a binding agent that binds to the proteins of the invention and determining the amount of bound protein in the sample. The polypeptides may be used as antigens in the production of antibodies pecific for P. acnes proteins. These antibodies can be used to downregulate expression and activity of P. acnes polypeptides and therefore treat P. acnes infections. The antibodies may also be used as diagnostic agents for determining P. acnes presence, for example, by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               enzyme linked immunosorbent assay (ELISA).
Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
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Pred. No. 35;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 62.1%;
Best Local Similarity 87.5%;
Matches 7; Conservative
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ID AAM9
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AC AAM9
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AAU48931 standard; Protein; 92 AA.

RESULT 11 AAU48931 AAU48931

1 NTLKTPRVGG 10 :|: |||:|| 2 HTIPTPRMGG 11

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us-09-743-225-8.rag

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The invention relates to primers for synthesising full length cDNA clones. 830 cDNA molecules encoding a human protein have been isolated and nucleotide sequences of 5'- and 3'-ends of the cDNA molecules have been determined. Primers for synthesising the full length cDNA are useful for clarifying the function of the protein encoded by the cDNA. The full length clones were obtained by construction of full length enriched cDNA libraries that were synthesised by the oligo-capping method. The primers enable the production of the full length cDNA encoded by a full length human cDNA of the invention.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in CD-ROM format directly from EPO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      830 Primers useful for synthesizing full length cDNA clones and their
                                                                                                                                                                                                                                                                                                                                                                                                                                       Ishii S, Kawai Y;
S, Otsuki T, Koga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Biosynthesis; biodegradation; lactic bacterium; yogurt; cheese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 8; SEQ ID NO 3731; 1380pp + sequence listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22; Length 140
                                                                               Human; full length cDNA; cDNA synthesis; oligo-capping.
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Wakamatsu A, Sugiyama T, Nagai K, Kojima S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 36; DB 2
Pred. No. 54;
2; Mismatches
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                                      Human polypeptide, SEQ ID NO: 3731
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lactococcus lactis protein ylfH.
                                                                                                                                                                                                                                                                                                   ; 2000JP-0194486.
; 2000JP-0118774.
; 2000JP-0183765.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               62.1%;
58.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             use in genetic manipulation
                                                                                                                                                                                                                                                            07-JUL-2000; 2000EP-0114089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 58.35,
Acceptage 7; Conservative
(first entry)
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N-PSDB; AAK94702.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      140 AA;
                                                                                                                                                                                                                                                                                                   08-JUL-1999;
11-JAN-2000;
02-MAY-2000;
                                                                                                                               Homo sapiens.
                                                                                                                                                                        EP1130094-A2.
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06-NOV-2001
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sequence (ABA90521) and related proteins (ABB53300-ABB55621). The nucleic acid sequence is useful in the detection and/or amplification of nucleic acid sequence, particularly to identify Lactococcus lactis or related species. The proteins of lanetion are useful for the biosynthesis or biodegradation of a composition of interest. The invention helps research in lactic bacteria, particularly useful in the production of yourt and cheese.

Note: The sequence data for this patent is based on equivalent patent WO200177334 (published 18-OCT-2001) which is available in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Protein identification; signal transduction pathway; metabolic pathway; hybridisation assay; genetic mapping; gene expression control; promoter; termination sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                               New nucleotide sequence useful in the identification or Lactococcus lactis and related species \cdot
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                                                                                                                                                                                                                                                                                                                                                                                                               Score 36; DB 23; Length 256;
Pred'. No. 1e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                      present invention is related to a Lactococcus lactis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Arabidopsis thaliana protein fragment SEQ ID NO: 3150
                                                                                            Renault P, Ehrlich SD;
                                                                                                                                                                                         Claim 6; SEQ ID No 1162; 2504pp; French.
                                                                 (INRG ) INRA INST NAT RECH AGRONOMIQUE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAG06393 standard; Protein; 319 AA.
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87.5%;
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99US-0128234.
99US-0128714.
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990S-0125788
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99US-0130449
                                     11-APR-2000; 2000FR-0004630
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           11-APR-2000; 2000FR-0004630
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Best Local Similarity 87.5
Local 7; Conservative
                                                                                            Bolotine A, Sorokine A,
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47 NTTKTPRV 54
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                                                                                                                      WPI; 2002-043418/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 NTLKTPRV 8
                                                                                                                                                                                                                                                                                                                                                                                         256 AA;
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21-APR-1999
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90S-0130510 90S-0130810 90S-0131449 90S-0132407 90S-0132485 90S-0132485 90S-0132485 90S-0132485 90S-0132485 90S-0132485	905 - 0134370 905 - 0134370 905 - 0135124 905 - 0135123 905 - 013523 905 - 013523 905 - 0137528 905 - 0137528 905 - 0137528 905 - 0137528 905 - 0137750 905 - 0139453 905 - 0139455 905 - 0139455 905 - 0139455 905 - 0139455 905 - 0139455 905 - 0139455	9005-0139463 9005-0139463 9005-0139463 9005-0139817. 9005-0139819. 9005-0140354. 9005-0140695. 9005-0141287. 9005-0141287. 9005-0141287. 9005-0141287. 9005-0141287. 9005-014205. 9005-014205. 9005-014205. 9005-014333. 9005-0144333. 9005-0144333. 9005-0144333. 9005-0144333. 9005-0144333.
3. APR-1999 8. APR-1999 8. APR-1999 0. APR-1999 0. APR-1999 6. MAX-1999 6. MAX-1999 7. MAY-1999 4. MAY-1999	4 - MAY - 1999 9 B MAY - 1999 9 B MAY - 1999 10 MAY - 1999 11 MAY - 1999 12 MAY - 1999 13 MAY - 1999 14 MAY - 1999 15 MAY - 1999 16 MAY - 1999 17 MAY - 1999 18 MAY - 1999 18 MAY - 1999 19 MAY - 1999	18-70N-1999; 18-70N-1999; 21-70N-1999; 22-70N-1999; 23-70N-1999; 24-70N-1999; 24-70N-1999; 24-70N-1999; 24-70N-1999; 10-70L-1999; 10-70L-1999; 11-70L-1999;
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PR 23-JUL-1999; 990S-0145148.
PR 23-JUL-1999; 990S-014518.
PR 23-JUL-1999; 990S-01518.
PR 23-JUL-19

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                                                                                                                                        Gaps
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                                                                                                                     Length 319;
                                                                                                                    Score 36; DB 21; Length 31
Pred. No. 1.38+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                              Arabidopsis thaliana protein fragment SEQ ID NO: 3149.
                                                                                                                                                                                                                        AAG06392 standard; Protein; 323 AA
                                                                                                                                                                                                                                                                                                                                                                                                           990S-0121825.
990S-0123180.
990S-0123548.
990S-0125788.
990S-0126264.
                                                                                                                      Query Match
Best Local Similarity 77.8%;
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99US-0128234.
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99US-0160989.
99US-0161404.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-0130077
99US-0130449
                                                                                                                                                                                                                                                                                                                                                                                         25-FEB-2000; 2000EP-0301439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-0127462
                                                                9US-016136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-013248
                                                                                                                                                                                                                                                            17-OCT-2000 (first entry)
                                                                                                                                                         2 TLKTPRVGG 10
                                                                                                                                                                   ||| | | |||
60 TLKVPTVGG 68
                                                                                                                                                                                                                                                                                                                                   Arabidopsis thaliana
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PR 22-JUL-1999, 9905-0145182

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PR 25-MC-1999, 9905-015436

PR 24-SEP-1999, 9905-015436

PR 24-SEP
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PR 21-0CT-1999; 9908-0160815.
PR 22-0CT-1999; 9908-0160881.
PR 22-0CT-1999; 9908-0160881.
PR 22-0CT-1999; 9908-0160881.
PR 22-0CT-1999; 9908-0161004.
PR 25-0CT-1999; 9908-0161004.
PR 25-0CT-1999; 9908-0161135.
PR 26-0CT-1999; 9908-0161136.
PR 26-0CT-1999; 9908-0161136.
PR 28-0CT-1999; 9908-0161136.
PR 28-0CT-1999; 9908-016192.
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PR 28-0CT-1999; 9908-016193.
PR 29-0CT-1999; 9008-0162142.

Query Match 62.1%; Score 36; DB 21; Length 323; Best Local Similarity 77.8%; Pred. No. 1.3e+02; Matches 7; Conservative 0; Mismatches 2; Indels 0; Gaps Qy 2 TiKTPRVGG 10

Qy 2 TiKTPRVGG 10

Gy 2 TiKTPRVGG 72

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Job time: 38.0909 secs